

Library preparation and sequencing

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An abbreviated version of this protocol was published in eLIFE in Aug 2020

Single-cell RNA-sequencing reveals distinct patterns of cell state heterogeneity in mouse models of breast cancer

DOI: [10.7554/eLife.58810](https://doi.org/10.7554/eLife.58810)

Related files



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How to cite: (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Yeo, S. and Guan, J. (2022). Library preparation and sequencing. Bio-protocol Preprint. bio-protocol.org/prep1525.
2. Yeo, S. K., Zhu, X., Okamoto, T., Hao, M., Wang, C., Lu, P., Lu, L. J. and Guan, J. (2020). Single-cell RNA-sequencing reveals distinct patterns of cell state heterogeneity in mouse models of breast cancer. eLIFE. DOI: [10.7554/eLife.58810](https://doi.org/10.7554/eLife.58810)

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